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Cys Arg Thr Ile Pro Glu Ala Cys Arg Gly Asp Met Met Cys Val 50 55 60

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Phe	Thr	Leu	Asn	Glu 200	Asp	Gly	Arg	Ser	Cys 205	Gln	Asp	Val	Asn	Glu 210
Cys	Ala	Thr	Glu	Asn 215	Pro	Cys	Val	Gln	Thr 220	Cys	Val	Asn	Thr	Tyr 225
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Phe	Leu	Cys	Gln	His 260	Glu	Cys	Val	Asn	Gln 265	Pro	Gly	Thr	Tyr	Phe 270
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Arg	_	Met											Gln	
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Met Thr Arg Pro	Ile Lys Gly 410	Pro Arg Glu Ile Gln 415	Leu Asp Leu 420
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- Pro Gly Phe Pro Ala Asn Val Thr Thr Leu Ser Leu Ser Ala Asn 50 55 60
- Arg Leu Pro Gly Leu Pro Glu Gly Ala Phe Arg Glu Val Pro Leu 65 70 75
- Leu Gln Ser Leu Trp Leu Ala His Asn Glu Ile Arg Thr Val Ala 80 85 90
- Ala Gly Ala Leu Ala Ser Leu Ser His Leu Lys Ser Leu Asp Leu
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- Ser His Asn Leu Ile Ser Asp Phe Ala Trp Ser Asp Leu His Asn 110 115 120
- Leu Ser Ala Leu Gln Leu Leu Lys Met Asp Ser Asn Glu Leu Thr
 125 130 135
- Phe Ile Pro Arg Asp Ala Phe Arg Ser Leu Arg Ala Leu Arg Ser 140 145 150
- Leu Gln Leu Asn His Asn Arg Leu His Thr Leu Ala Glu Gly Thr
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- Phe Thr Pro Leu Thr Ala Leu Ser His Leu Gln Ile Asn Glu Asn 170 175 180
- Pro Phe Asp Cys Thr Cys Gly Ile Val Trp Leu Lys Thr Trp Ala 185 190 195
- Leu Thr Thr Ala Val Ser Ile Pro Glu Gln Asp Asn Ile Ala Cys 200 205 210
- Thr Ser Pro His Val Leu Lys Gly Thr Pro Leu Ser Arg Leu Pro 215 220 225

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Asp Phe Gly Lys Leu Glu Glu Gly Thr Tyr Ser Cys Leu Ala Thr
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Glu	Val	Asp	Val	Ile 155	Val	Met	Asn	Ser	Glu 160	Gly	Asn	Thr	Ile	Leu 165
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Lys Thr Cys Cys Phe Gln Tyr Ser His Lys Pro Leu Pro Trp Thr
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 Trp Val Arg Ser Tyr Glu Phe Thr Ser Asn Ser Cys Ser Gln Arg
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Ala Val Ile Phe Thr Thr Lys Arg Gly Lys Lys Val Cys Thr His
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 Pro Lys Gln Leu
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<223> Synthetic oligonucleotide probe
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<400> 27
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<211> 3552
<212> DNA
<213> Homo Sapien
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<211> 386

<212> PRT

<213> Homo Sapien

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35 40 45

Ala Val Leu Leu Pro Val Arg Val Asp Ser Ala Thr Ile Pro Arg
50 55 60

Gln Asp Glu Val Pro Gln Gln Thr Val Ala Pro Gln Gln Gln Arg
65 70 75

Arg Ser Leu Lys Glu Glu Glu Cys Pro Ala Gly Ser His Arg Ser 80 85 90

Glu Tyr Thr Gly Ala Cys Asn Pro Cys Thr Glu Gly Val Asp Tyr 95 100 105

Thr Ile Ala Ser Asn Asn Leu Pro Ser Cys Leu Leu Cys Thr Val

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Asp	Thr	Val	Cys	Gln 140	Cys	Glu	Lys	Gly	Ser 145	Phe	Gln	Asp	Lys	Asn 150
Ser	Pro	Glu	Met	Cys 155	Arg	Thr	Cys	Arg	Thr 160	Gly	Cys	Pro	Arg	Gly 165
Met	Val	Lys	Val	Ser 170	Asn	Cys	Thr	Pro	Arg 175	Ser	Asp	Ile	Lys	Cys 180
Lys	Asn	Glu	Ser	Ala 185	Ala	Ser	Ser	Thr	Gly 190	Lys	Thr	Pro	Ala	Ala 195
Glu	Glu	Thr	Val	Thr 200	Thr	Ile	Leu	Gly	Met 205	Leu	Ala	Ser	Pro	Tyr 210
His	Tyr	Leu	Ile	Ile 215	Ile	Val	Val	Leu	Val 220	Ile	Ile	Leu	Ala	Val 225
Val	Val	Val	Gly	Phe 230	Ser	Cys	Arg	Lys	Lys 235	Phe	Ile	Ser	Tyr	Leu 240
Lys	Gly	Ile	Cys	Ser 245	Gly	Gly	Gly	Gly	Gly 250	Pro	Glu	Arg	Val	His 255
Arg	Val	Leu	Phe	Arg 260	Arg	Arg	Ser	Cys	Pro 265	Ser	Arg	Val	Pro	Gly 270
Ala	Glu	Asp	Asn	Ala 275	Arg	Asn	Glu	Thr	Leu 280	Ser	Asn	Arg	Tyr	Leu 285
Gln	Pro	Thr	Gln	Val 290	Ser	Glu	Gln	Glu	Ile 295	Gln	Gly	Gln	Glu	Leu 300
Ala	Glu	Leu	Thr	Gly 305	Val	Thr	Val	Glu	Ser 310	Pro	Glu	Glu	Pro	Gln 315
Arg	Leu	Leu	Glu	Gln 320	Ala	Glu	Ala	Glu	Gly 325	Сув	Gln	Arg	Arg	Arg 330
Leu	Leu	Val	Pro	Val 335	Asn	Asp	Ala	Asp	Ser 340	Ala	Asp	Ile	Ser	Thr 345
Leu	Leu	Asp	Ala	Ser 350	Ala	Thr	Leu	Glu	Glu 355	Gly	His	Ala	Lys	Glu 360
Thr	Ile	Gln	Asp				Gly				Leu			Glu 375
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<213> Artificial Sequence

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agageteatt ceagatgeac ecetgteeag tgetgeetat ageateegea 150
gcatcgggga gaggcctgtc ctcaaagctc cagtccccaa aaggcaaaaa 200
 tgtgaccact ggactccctg cccatctgac acctatgcct acaggttact 250
 cagcggaggt ggcagaagca agtacgccaa aatctgcttt gaggataacc 300
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gtatgaaggc gataactctg gaccgatgac aaagtttatt cagagtgctg 450
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 caggaacatg aaattcaggt ctagctgggt atttattgca gcaaaaggct 600
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 aagaacaaca gatattctgg ctggcctgca gagatccaga tagaaggctg 700
 catacccaaa gaacgaagct gacactgcag ggtcctgagt aaatgtgttc 750
 tgtataaaca aatgcagctg gaatcgctca agaatcttat ttttctaaat 800
 ccaacagccc atatttgatg agtattttgg gtttgttgta aaccaatgaa 850
 catttgctag ttgtatcaaa tcttggtacg cagtattttt ataccagtat 900
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 cttaaaaaaa aaa 963
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<212> PRT
<213> Homo Sapien
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<400> 32

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Leu Ile Pro Asp Ala Pro Leu Ser Ser Ala Ala Tyr Ser Ile Arg
                  35
Ser Ile Gly Glu Arg Pro Val Leu Lys Ala Pro Val Pro Lys Arg
                                                           60
                  50
                                      55
Gln Lys Cys Asp His Trp Thr Pro Cys Pro Ser Asp Thr Tyr Ala
                                                           75
                  65
Tyr Arg Leu Leu Ser Gly Gly Gly Arg Ser Lys Tyr Ala Lys Ile
                                                           90
Cys Phe Glu Asp Asn Leu Leu Met Gly Glu Gln Leu Gly Asn Val
                                                          105
                  95
                                     100
Ala Arg Gly Ile Asn Ile Ala Ile Val Asn Tyr Val Thr Gly Asn
                                                          120
                                     115
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Val Thr Ala Thr Arg Cys Phe Asp Met Tyr Glu Gly Asp Asn Ser
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                                     130
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Gly Pro Met Thr Lys Phe Ile Gln Ser Ala Ala Pro Lys Ser Leu
                                                          150
                                     145
                 140
Leu Phe Met Val Thr Tyr Asp Asp Gly Ser Thr Arg Leu Asn Asn
                                     160
                                                          165
                 155
Asp Ala Lys Asn Ala Ile Glu Ala Leu Gly Ser Lys Glu Ile Arg
                                     175
                                                          180
                 170
Asn Met Lys Phe Arg Ser Ser Trp Val Phe Ile Ala Ala Lys Gly
                                                          195
                                     190
                 185
Leu Glu Leu Pro Ser Glu Ile Gln Arg Glu Lys Ile Asn His Ser
                           205
Asp Ala Lys Asn Asn Arg Tyr Ser Gly Trp Pro Ala Glu Ile Gln
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 Ile Glu Gly Cys Ile Pro Lys Glu Arg Ser
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<220>

<223> Synthetic oligonucleotide probe

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<213> Artificial Sequence
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<400> 35
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<211> 24
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<223> Synthetic oligonucleotide probe
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<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide probe
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<211> 1215
<212> DNA
<213> Homo Sapien
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 gctccaccgg ggtcgctcgc cctccggctc ctgctgttcg tggcgctacc 200
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- <210> 39
- <211> 330
- <212> PRT
- <213> Homo Sapien
- <400> 39
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- Phe Val Ala Leu Pro Ala Ser Gly Trp Leu Thr Thr Gly Ala Pro 20 25 30
- Glu Pro Pro Pro Leu Ser Gly Ala Pro Gln Asp Gly Ile Arg Ile
 35 40 45

Asn Val Thr Thr Leu Lys Asp Asp Gly Asp Ile Ser Lys Gln Gln

				50					55					60
Val	Val	Leu	Asn	Ile 65	Thr	Tyr	Glu	Ser	Gly 70	Gln	Val	Tyr	Val	Asn 75
Asp	Leu	Pro	Val	Asn 80	Ser	Gly	Val	Thr	Arg 85	Ile	Ser	Cys	Gln	Thr 90
Leu	Ile	Val	Lys	Asn 95	Glu	Asn	Leu	Glu	Asn 100	Leu	Glu	Glu	Lys	Glu 105
Tyr	Phe	Gly	Ile	Val 110	Ser	Val	Arg	Ile	Leu 115	Val	His	Glu	Trp	Pro 120
Met	Thr	Ser	Gly	Ser 125	Ser	Leu	Gln	Leu	Ile 130	Val	Ile	Gln	Glu	Glu 135
Val	Val	Glu	Ile	Asp 140	Gly	Lys	Gln	Val	Gln 145	Gln	Lys	Asp	Val	Thr 150
Glu	Ile	Asp	Ile	Leu 155	Val	Lys	Asn	Arg	Gly 160	Val	Leu	Arg	His	Ser 165
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Arg	Asp	Ser	Asp	Ile 185	Leu	Phe	Thr	Leu	Pro 190	Asn	Leu	Ser	Lys	Lys 195
Glu	Ser	Val	Ser	Ser 200	Leu	Gln	Thr	Thr	Ser 205	Gln	Tyr	Leu	Ile	Arg 210
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Pro	Glu	Thr	Pro	Leu 230	Arg	Ala	Glu	Pro	Pro 235	Ser	Ser	Tyr	Lys	Val 240
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Trp	Ser	Asn	Val	Phe 260	Pro	Val	Phe	Phe	Gln 265	Phe	Leu	Asn	Ile	Met 270
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Val	Phe	Phe	Pro	Val 290	Ser	Glu	Tyr	Lys	Gly 295	Ile	Leu	Gln	Leu	Asp 300
Lys	Val	Asp	Val	Ile 305	Pro	Val	Thr	Ala	Ile 310	Asn	Leu	Tyr	Pro	Asp 315
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<210> 40

<211> 2498

<212> DNA <213> Homo Sapien

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<211> 263

<212> PRT

<213> Homo Sapien

<400> 41

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Cys Cys Ala Ser Gly Asn Ile Asp Thr Ala Trp Ser Asn Leu Thr
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                 65
His Pro Gly Asn Lys Ile Asn Leu Leu Gly Phe Leu Gly Leu Val
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                                     85
                 80
His Cys Leu Pro Cys Lys Asp Ser Cys Asp Gly Val Glu Cys Gly
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Pro Gly Lys Ala Cys Arg Met Leu Gly Gly Arg Pro Arg Cys Glu
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Cys Ala Pro Asp Cys Ser Gly Leu Pro Ala Arg Leu Gln Val Cys
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                                    130
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Gly Ser Asp Gly Ala Thr Tyr Arg Asp Glu Cys Glu Leu Arg Ala
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                                    145
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Ala Arg Cys Arg Gly His Pro Asp Leu Ser Val Met Tyr Arg Gly
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                                    160
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Arg Cys Arg Lys Ser Cys Glu His Val Val Cys Pro Arg Pro Gln
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                                    175
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Ser Cys Val Val Asp Gln Thr Gly Ser Ala His Cys Val Val Cys
                                                        195
                                    190
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Arg Ala Ala Pro Cys Pro Val Pro Ser Ser Pro Gly Gln Glu Leu
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                200
Cys Gly Asn Asn Asn Val Thr Tyr Ile Ser Ser Cys His Met Arg
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Gln Ala Thr Cys Phe Leu Gly Arg Ser Ile Gly Val Arg His Ala
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Gly Ser Cys Ala Gly Thr Pro Glu Glu Pro Pro Gly Gly Glu Ser
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<220>

<223> Synthetic oligonucleotide probe

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 <223> Synthetic oligonucleotide probe
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 <212> DNA
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<212> DNA
<213> Artificial Sequence
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<211> 1690
<212> DNA
<213> Homo Sapien
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<211> 505

<212> PRT

<213> Homo Sapien

<400> 52

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Arg	Ala	Pro	Pro	Asp 35	Lys	Ile	Ala	Ile	Ile 40	Gly	Ala	Gly	Ile	G1y 45
Gly	Thr	Ser	Ala	Ala 50	Tyr	Tyr	Leu	Arg	Gln 55	Lys	Phe	Gly	Lys	Asp 60
Val	Lys	Ile	Asp	Leu 65	Phe	Glu	Arg	Glu	Glu 70	Val	Gly	Gly	Arg	Leu 75
Ala	Thr	Met	Met	Val 80	Gln	Gly	Gln	Glu	Tyr 85	Glu	Ala	Gly	Gly	Ser 90
Val	Ile	His	Pro	Leu 95	Asn	Leu	His	Met	Lys 100	Arg	Phe	Val	Lys	Asp 105
Leu	Gly	Leu	Ser	Ala 110	Val	Gln	Ala	Ser	Gly 115	Gly	Leu	Leu	Gly	Ile 120
Tyr	Asn	Gly	Glu	Thr 125	Leu	Val	Phe	Glu	Glu 130	Ser	Asn	Trp	Phe	Ile 135
Ile	Asn	Val	Ile	Lys 140	Leu	Val	Trp	Arg	Tyr 145	Gly	Phe	Gln	Ser	Leu 150
Arg	Met	His	Met	Trp 155	Val	Glu	Asp	Val	Leu 160	Asp	Lys	Phe	Met	Arg 165
Ile	Tyr	Arg	Tyr	Gln 170	Ser	His	Asp	Tyr	Ala 175	Phe	Ser	Ser	Val	Glu 180
Lys	Leu	Leu	His	Ala 185	Leu	Gly	Gly	Asp	Asp 190	Phe	Leu	Gly	Met	Leu 195
Asn	Arg	Thr	Leu	Leu 200	Glu	Thr	Leu	Gln	Lys 205	Ala	Gly	Phe	Ser	Glu 210
Lys	Phe	Leu	Asn	Glu 215	Met	Ile	Ala	Pro	Val 220	Met	Arg	Val	Asn	Tyr 225
Gly	Gln	Ser	Thr	Asp 230	Ile	Asn	Ala	Phe	Val 235	Gly	Ala	Val	Ser	Leu 240
Ser	Cys	Ser	Asp	Ser 245	Gly	Leu	Trp	Ala	Val 250	Glu	Gly	Gly	Asn	Lys 255
Leu	Val	Cys	Ser	Gly 260		Leu	Gln	Ala	Ser 265		Ser	Asn	Leu	Ile 270
Ser	Gly	Ser	Val	Met 275		Ile	Glu	Glu	Lys 280	Thr	Lys	Thr	Lys	Tyr 285
Thr	Gly	Asn	Pro	Thr 290		Met	Tyr	Glu	Val 295	Val	Tyr	Gln	Ile	Gly 300
Thr	Glu	Thr	Arg	Ser 305			. Tyr		Ile 310	Val	Leu	val	Ala	Thr 315
Pro	Leu	Asn	Arq	Lys	Met	Ser	Asn	Ile	Thr	Phe	Leu	. Asn	Phe	Asp

				320					325					330
Pro	Pro	Ile	Glu	Glu 335	Phe	His	Gln	Tyr	Tyr 340	Gln	His	Ile	Val	Thr 345
Thr	Leu	Val	Lys	Gly 350	Glu	Leu	Asn	Thr	Ser 355	Ile	Phe	Ser	Ser	Arg 360
Pro	Ile	Asp	Lys	Phe 365	Gly	Leu	Asn	Thr	Val 370	Leu	Thr	Thr	Asp	Asn 375
Ser	Asp	Leu	Phe	Ile 380	Asn	Ser	Ile	Gly	Ile 385	Val	Pro	Ser	Val	Arg 390
Glu	Lys	Glu	Asp	Pro 395	Glu	Pro	Ser	Thr	Asp 400	Gly	Thr	Tyr	Val	Trp 405
Lys	Ile	Phe	Ser	Gln 410	Glu	Thr	Leu	Thr	Lys 415	Ala	Gln	Ile	Leu	Lys 420
Leu	Phe	Leu	Ser	Tyr 425	Asp	Tyr	Ala	Val	Lys 430	Lys	Pro	Trp	Leu	Ala 435
Tyr	Pro	His	Tyr	Lys 440	Pro	Pro	Glu	Lys	Cys 445	Pro	Ser	Ile	Ile	Leu 450
His	Asp	Arg	Leu	Tyr 455	Tyr	Leu	Asn	Gly	Ile 460	Glu	Cys	Ala	Ala	Ser 465
Ala	Met	Glu	Met	Ser 470	Ala	Ile	Ala	Ala	His 475	Asn	Ala	Ala	Leu	Leu 480
Ala	Tyr	His	Arg	Trp 485	Asn	Gly	His	Thr	Asp 490	Met	Ile	Asp	Gln	Asp 495
Gly	Leu	Tyr	Glu	Lys 500	Leu	Lys	Thr	Glu	Leu 505					

<210> 53

<211> 728

<212> DNA

<213> Homo Sapien

<400> 53

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aaaccttatg catgaaacca ctgataagaa tttatcacct gatgggcaat 400 atgtgcctag aatcatgttt gtagaccctt ctttaacagt tagagctgac 450 atagctggaa gatactctaa cagattgtac acatatgagc ctcgggattt 500 acccctattg atagaaaaca tgaagaaagc attaagactt attcagtcag 550 agctataaga gatgatggaa aaaagccttc acttcaaaga agtcaaattt 600 catgaagaaa acctctggca cattgacaaa tactaaatgt gcaagtatat 650 agattttgta atattactat ttagtttttt taatgtgttt gcaatagtct 700 tattaaaata aatgtttttt aaatctga 728

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155

160

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<211> 537
<212> DNA
<213> Homo Sapien
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 agaccaggaa cgagaaaaaa gaagtatcag tgacagcgat gaattagctt 200
 cagggttttt tgtgttccct tacccatatc catttcgccc acttccacca 250
 attccatttc caagatttcc atggtttaga cgtaattttc ctattccaat 300
 acctgaatct gcccctacaa ctccccttcc tagcgaaaag taaacaagaa 350
 ggataagtca cgataaacct ggtcacctga aattgaaatt gagccacttc 400
 cttgaagaat caaaattcct gttaataaaa gaaaaacaaa tgtaattgaa 450
 atagcacaca gcattctcta gtcaatatct ttagtgatct tctttaataa 500
 acatgaaagc aaagattttg gtttcttaat ttccaca 537
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<211> 85
<212> PRT
<213> Homo Sapien
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                                                           30
                                       25
                  20
 Ile Ser Asp Ser Asp Glu Leu Ala Ser Gly Phe Phe Val Phe Pro
 Tyr Pro Tyr Pro Phe Arg Pro Leu Pro Pro Ile Pro Phe Pro Arg
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                  50
 Phe Pro Trp Phe Arg Arg Asn Phe Pro Ile Pro Ile Pro Glu Ser
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 Ala Pro Thr Thr Pro Leu Pro Ser Glu Lys
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<210> 57
<211> 2997
<212> DNA
<213> Homo Sapien - - -
<400> 57
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- <210> 58 <211> 747
- <212> PRT

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- <213> Homo Sapien
- <400> 58
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10

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- Ala Thr Cys Gln Gln Arg Glu Gly Lys Lys Ile Cys Ile Cys Asn 50 55 60
- Tyr Gly Phe Val Gly Asn Gly Arg Thr Gln Cys Val Asp Lys Asn 65 70 75
- Glu Cys Gln Phe Gly Ala Thr Leu Val Cys Gly Asn His Thr Ser 80 85 90
- Cys His Asn Thr Pro Gly Gly Phe Tyr Cys Ile Cys Leu Glu Gly 95 100 105
- Tyr Arg Ala Thr Asn Asn Asn Lys Thr Phe Ile Pro Asn Asp Gly 110 115 120
- Thr Phe Cys Thr Asp Ile Asp Glu Cys Glu Val Ser Gly Leu Cys 125 130 135
- Arg His Gly Gly Arg Cys Val Asn Thr His Gly Ser Phe Glu Cys . 140 145 150
- Tyr Cys Met Asp Gly Tyr Leu Pro Arg Asn Gly Pro Glu Pro Phe 155 160 165
- His Pro Thr Thr Asp Ala Thr Ser Cys Thr Glu Ile Asp Cys Gly 170 175 180
- Thr Pro Pro Glu Val Pro Asp Gly Tyr Ile Ile Gly Asn Tyr Thr 185 190 195
- Ser Ser Leu Gly Ser Gln Val Arg Tyr Ala Cys Arg Glu Gly Phe 200 205 210
- Phe Ser Val Pro Glu Asp Thr Val Ser Ser Cys Thr Gly Leu Gly 215
- Thr Trp Glu Ser Pro Lys Leu His Cys Gln Glu Ile Asn Cys Gly 230 235 240

Asn	Pro	Pro	GIU	Met 245	arg	HIS	АТА	116	250	vai	GIÀ	ASI	HIS	255
Ser	Arg	Leu	Gly	Gly 260	Val	Ala	Arg	Tyr	Val 265	Cys	Gln	Glu	Gly	Phe 270
Glu	Ser	Pro	Gly	Gly 275	Lys	Ile	Thr	Ser	Val 280	Cys	Thr	Glu	Lys	Gly 285
Thr	Trp	Arg	Glu	Ser 290	Thr	Leu	Thr	CAa	Thr 295	Glu	Ile	Leu	Thr	100
Ile	Asn	Asp	Val	Ser 305	Leu	Phe	Asn	Asp	Thr 310	Cys	Val	Arg	Trp	Gln 315
Ile	Asn	Ser	Arg	Arg 320	Ile	Asn	Pro	Lys	Ile 325	Ser	Tyr	Val	Ile	Ser 330
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Thr	Val	Asn	Leu	Thr 350	Thr	Asp	Ser	Arg	Thr 355	Pro	Glu	Val	Cys	Leu 360
Ala	Leu	Tyr	Pro	Gly 365	Thr	Asn	Tyr	Thr	Val 370	Asn	Ile	Ser	Thr	Ala 375
Pro	Pro	Arg	Arg	Ser 380	Met	Pro	Ala	Val	Ile 3,85	Gly	Phe	Gln	Thr	Ala 390
Glu	Val	Asp	Leu	Leu 395	Glu	Asp	Asp	Gly	Ser 400	Phe	Asn	Ile	Ser	Ile 405
Phe	Asn	Glu	Thr	Cys 410	Leu	Lys	Leu	Asn	Arg 415	Arg	Ser	Arg	Lys	Val 420
Gly	Ser	Glu	His	Met 425	Tyr	Gln	Phe	Thr	Val 430	Leu	Gly	Gln	Arg	Trp 435
Tyr	Leu	Ala	Asn	Phe 440	Ser	His	Ala	Thr	Ser 445	Phe	Asn	Phe	Thr	Thr 450
Arg	Glu	Gln	Val	Pro 455	Val	Val	Cys	Leu	Asp 460	Leu	Tyr	Pro	Thr	Thr 465
Asp	Tyr	Thr	Val	Asn 470	Val	Thr	Leu	Leu	Arg 475	Ser	Pro	Lys	Arg	His 480
Ser	Val	Gln	Ile	Thr 485	Ile	Ala	Thr	Pro	Pro 490	Ala	Val	Lys	Gln	Thr 495
Ile	Ser	Asn	Ile	Ser 500	Gly	Phe	Asn	Glu	Thr 505	Cys	Leu	Arg	Trp	Arg 510
Ser	Ile	Lys		Ala _515	_		Glu	Glu	Met 520	Tyr	Leu	Phe	His	Ile 525
Trp	Gly	Gln	Arq	Trp	Tyr	Gln	Lys	Glu	Phe	Ala	Gln	Glu	Met	Thr

				530					535					540
Phe	Asn	Ile	Ser	Ser 545	Ser	Ser	Arg	Asp	Pro 550	Glu	Val	Cys	Leu	Asp 555
Leu	Arg	Pro	Gly	Thr 560	Asn	Tyr	Asn	Val	Ser 565	Leu	Arg	Ala	Leu	Ser 570
Ser	Glu	Leu	Pro	Val 575	Val	Ile	Ser	Leu	Thr 580	Thr	Gln	Ile	Thr	Glu 585
Pro	Pro	Leu	Pro	Glu 590	Val	Glu	Phe	Phe	Thr 595	Val	His	Arg	Gly	Pro 600
Leu	Pro	Arg	Leu	Arg 605	Leu	Arg	Lys	Ala	Lys	Glu	Lys	Asn	Gly	Pro 615
Ile	Ser	Ser	Tyr	Gln 620	Val	Leu	Val	Leu	Pro 625	Leu	Ala	Leu	Gln	Ser 630
Thr	Phe	Ser	Cys	Asp 635	Ser	Glu	Gly	Ala	Ser 640	Ser	Phe	Phe	Ser	Asn 645
Ala	Ser	Asp	Ala	Asp 650	Gly	Tyr	Val	Ala	Ala 655	Glu	Leu	Leu	Ala	Lys 660
Asp	Val	Pro	Asp	Asp 665	Ala	Met	Glu	Ile	Pro 670	Ile	Gly	Asp	Arg	Leu 675
Tyr	Tyr	Gly	Glu	Tyr 680	Tyr	Asn	Ala	Pro	Leu 685	Lys	Arg	Gly	Ser	Asp 690
Tyr	Cys	Ile	Ile	Leu 695	Arg	Ile	Thr	Ser	Glu 700	Trp	Asn	Lys	Val	Arg 705
Arg	His	Ser	Cys	Ala 710	Val	Trp	Ala	Gln	Val 715	Lys	Asp	Ser	Ser	Leu 720
Met	Leu	Leu	Gln	Met 725	Ala	Gly	Val	Gly	Leu 730	Gly	Ser	Leu	Ala	Val 735
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<211> 482

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<213> Homo Sapien

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Ser	Ala	Glu	Thr	Ser 65	Ser	Arg	Ala	Ser	Thr 70	Pro	Ala	Gly	Pro	Ile 75
Pro	Glu	Ala	Glu	Thr 80	Arg	Gly	Ala	Lys	Arg 85	Ile	Ser	Pro	Ala	Arg 90
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Ile	Ala	Thr	Ser	Val 110	Glu	Thr	Ser	Ala	Ala 115	Ser	Gly	Ser	Pro	Glu 120
Gly	Ala	Gly	Met	Thr 125	Thr	Val	Gln	Thr	Ile 130	Thr	Gly	Ser	Asp	Pro 135
Glu	Glu	Ala	Ile	Phe 140	Asp	Thr	Leu	Cys	Thr 145	Asp	Asp	Ser	Ser	Glu 150
Glu	Ala	Lys	Thr	Leu 155	Thr	Met	Asp	Ile	Leu 160	Thr	Leu	Ala	His	Thr 165
Ser	Thr	Glu	Ala	Lys 170	Gly	Leu	Ser	Ser	Glu 175	Ser	Ser	Ala	Ser	Ser 180
Asp	Gly	Pro	His	Pro 185	Val	Ile	Thr	Pro	Ser 190	Arg	Ala	Ser	Glu	Ser 195
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			Ser	215					220					225
			Trp	230					235					240
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Thr	Glu	Ile	Glu	Thr 260	Thr	Thr	Ser	Ser	Ile 265	Pro	Gly	Ala	Ser	Asp 270
Ile	Asp	Leu	Ile	Pro 275	Thr	Glu	Gly	Val	Lys 280	Ala	Ser	Ser	Thr	Ser 285
Asp	Pro	Pro	Ala	Leu 290	Pro	Asp	Ser	Thr	Glu 295	Ala	Lys	Pro	His	11e 300
Thr	Glu	Val	Thr	Ala 305	Ser	Ala	Glu	Thr	Leu 310	Ser	Thr	Ala	Gly	Thr 315

Thr	Glu	Ser	Ala	Ala 320	Pro	His	Ala	Thr	Val 325	Gly	Thr	Pro	Leu	Pro 330
Thr	Asn	Ser	Ala	Thr 335	Glu	Arg	Glu	Val	Thr 340	Ala	Pro	Gly	Ala	Thr 345
Thr	Leu	Ser	Gly	Ala 350	Leu	Val	Thr	Val	Ser 355	Arg	Asn	Pro	Leu	Glu 360
Glu	Thr	Ser	Ala	Leu 365	Ser	Val	Glu	Thr	Pro 370	Ser	Tyr	Val	Lys	Val 375
Ser	Gly	Ala	Ala	Pro 380	Val	Ser	Ile	Gľu	Ala 385	Gly	Ser	Ala	Val	Gly 390
Lys	Thr	Thr	Ser	Phe 395	Ala	Gly	Ser	Ser	Ala 400	Ser	Ser	Tyr	Ser	Pro 405
Ser	Glu	Ala	Ala	Leu 410	Lys	Asn	Phe	Thr	Pro 415	Ser	Glu	Thr	Pro	Thr 420
Met	Asp	Ile	Ala	Thr 425	Lys	Gly	Pro	Phe	Pro 430	Thr	Ser	Arg	Asp	Pro 435
Leu	Pro	Ser	Val	Pro 440	Pro	Thr	Thr	Thr	Asn 445	Ser	Ser	Arg	Gly	Thr 450
Asn	Ser	Thr	Leu	Ala 455	Lys	Ile	Thr	Thr	Ser 460	Ala	Lys	Thr	Thr	Met 465
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aagaaaattt cccaagcttc ctcctgcctg cagaagctcc tctactttaa 400

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- <211> 364
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- Gln Phe Leu Gly Leu Asp Lys Ala Pro Ser Pro Gln Lys Phe Gln 40
- Pro Val Pro Tyr Ile Leu Lys Lys Ile Phe Gln Asp Arg Glu Ala 60 55 50
- Ala Ala Thr Thr Gly Val Ser Arg Asp Leu Cys Tyr Val Lys Glu 75 -65 70
- Leu Gly Val Arg Gly Asn Val Leu Arg Phe Leu Pro Asp Gln Gly

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Gln	Lys	Leu	Leu	Tyr 110	Phe	Asn	Leu	Ser	Ala 115	Ile	Lys	Glu	Arg	Glu 120
Gln	Leu	Thr	Leu	Ala 125	Gln	Leu	Gly	Leu	Asp 130	Leu	Gly	Pro	Asn	Ser 135
Tyr	Tyr	Asn	Leu	Gly 140	Pro	Glu	Leu	Glu	Leu 145	Ala	Leu	Phe	Leu	Val 150
Gln	Glu	Pro	His	Val 155	Trp	Gly	Gln	Thr	Thr 160	Pro	Lys	Pro	Gly	Lys 165
Met	Phe	Val	Leu	Arg 170	Ser	Val	Pro	Trp	Pro 175	Gln	Gly	Ala	Val	His 180
Phe	Asn	Leu	Leu	Asp 185	Val	Ala	Lys	Asp	Trp 190	Asn	Asp	Asn	Pro	Arg 195
Lys	Asn	Phe	Gly	Leu 200	Phe	Leu	Glu	Ile	Leu 205	Val	Lys	Glu	Asp	Arg 210
Asp	Ser	Gly	Val	Asn 215	Phe	Gln	Pro	Glu	Asp 220	Thr	Cys	Ala	Arg	Leu 225
Arg	Cys	Ser	Leu	His 230	Ala	Ser	Leu	Leu	Val 235	Val	Thr	Leu	Asn	Pro 240
Asp	Gln	Cys	His	Pro 245	Ser	Arg	Lys	Arg	Arg 250	Ala	Ala	Ile	Pro	Val 255
Pro	Lys	Leu	Ser	Cys 260	Lys	Asn	Leu	Cys	His 265	Arg	His	Gln	Leu	Phe 270
Ile	Asn	Phe	Arg	Asp 275	Leu	Gly	Trp	His	Lys 280	Trp	Ile	Ile	Ala	Pro 285
Lys	Gly	Phe	Met	Ala 290	Asn	Tyr	Cys	His	Gly 295	Glu	Cys	Pro	Phe	Ser 300
Leu	Thr	Ile	Ser	Leu 305	Asn	Ser	Ser	Asn	Tyr 310	Ala	Phe	Met	Gln	Ala 315
Leu	Met	His	Ala	Val 320	Asp	Pro	Glu	Ile	Pro 325	Gln	Ala	Val	Cys	Ile 330
Pro	Thr	Lys	Leu	Ser 335	Pro	Ile	Ser	Met	Leu 340	Tyr	Gln	Asp	Asn	Asn 345
Asp	Asn	Val	Ile	Leu 350	Arg	His	Tyr	Glu	Asp 355	Met	Val	Val	Asp	Glu 360
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<400> 67
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<212> PRT

<213> Homo Sapien

<400> 73

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Gln Gly Glu Gln Glu Trp Glu Gly Thr Glu Glu Leu Pro Ser 35 40 45

Pro Pro Asp His Ala Glu Arg Ala Glu Glu Gln His Glu Lys Tyr
50 55 60

Arg Pro Ser Gln Asp Gln Gly Leu Pro Ala Ser Arg Cys Leu Arg
65 70 75

Cys Cys Asp Pro Gly Thr Ser Met Tyr Pro Ala Thr Ala Val Pro 80 85 90

Gln Ile Asn Ile Thr Ile Leu Lys Gly Glu Lys Gly Asp Arg Gly
95 100 105

Asp Arg Gly Leu Gln Gly Lys Tyr Gly Lys Thr Gly Ser Ala Gly 110 115 120

Ala Arg Gly His Thr Gly Pro Lys Gly Gln Lys Gly Ser Met Gly 125 130 135

Ala Pro Gly Glu Arg Cys Lys Ser His Tyr Ala Ala Phe Ser Val 140 145 150

Gly Arg Lys Lys Pro Met His Ser Asn His Tyr Tyr Gln Thr Val

Ile Phe Asp Thr Glu Phe Val Asn Leu Tyr Asp His Phe Asn Met 170 175 180

Phe Thr Gly Lys Phe Tyr Cys Tyr Val Pro Gly Leu Tyr Phe Phe 185 195

Ser Leu Asn Val His Thr Trp Asn Gln Lys Glu Thr Tyr Leu His

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<211> 167

<212> PRT

<213> Homo Sapien

<400> 78

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Arg Phe Leu Pro Glu Asp Leu Lys Gln Lys Glu Phe Pro Pro Ala
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                 95
                                     100
Met Lys Leu Leu Tyr Ser Val Glu His Glu Lys Pro Leu Tyr Leu
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                                     115
                110
Ser Phe Gly Arg Pro Glu Asn Lys Arg Ile Phe Pro Phe Pro Ile
                                                          135
                                     130
                125
Arg Glu Thr Ser Arg His Phe Ala Asp Leu Ala His Asn Ser Asp
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- <212> DNA
- <213> Homo Sapien
- <220>
- <221> unsure
- <222> 794
- <223> unknown base
- <400> 79
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 ggaagcagga accaagctta ggctgctcca tcccagctat ccttgttettg 200
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 cagcccaggg ctgcaggaag gacaggggg cctccaagac tggcaagaaa 350
 ggaaagggct ccaaaggctg caagaggact gagcggtcac agaccctaa 400
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gctttaacca ccccatctgc attcccagct ctaccctgca tggctgagct 600

geccacagea ggecaggtec agagagaceg aggaggaga gteteccagg 650 gageatgaga ggaggeagea ggaetgtece ettgaaggag aateateagg 700 accetggace tgataegget ecceagtaca ecceacetet teettgtaaa 750 tatgatttat acctaactga ataaaaaget gttetgtett eccneeca 798

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<211> 134

<212> PRT

<213> Homo Sapien

<400> 80

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Cys Cys Leu Lys Tyr Ser Gln Arg Lys Ile Pro Ala Lys Val Val
35 40 45

Arg Ser Tyr Arg Lys Gln Glu Pro Ser Leu Gly Cys Ser Ile Pro 50 55 60

Ala Ile Leu Phe Leu Pro Arg Lys Arg Ser Gln Ala Glu Leu Cys
65 70 75

Ala Asp Pro Lys Glu Leu Trp Val Gln Gln Leu Met Gln His Leu 80 85 90

Asp Lys Thr Pro Ser Pro Gln Lys Pro Ala Gln Gly Cys Arg Lys
95 100 105

Asp Arg Gly Ala Ser Lys Thr Gly Lys Lys Gly Lys Gly Ser Lys

Gly Cys Lys Arg Thr Glu Arg Ser Gln Thr Pro Lys Gly Pro 125 130

<210> 81

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Synthetic oligonucleotide probe

<400> 81

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<210> 82

<211> 19

<212> DNA

<213> Artificial Sequence

<220>

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<210> 83
<211> 924
<212> DNA
<213> Homo Sapien
<400> 83
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cggtctcagg agatgtctga tttccacaga catgcaccat atagaagaga 150
gtttccaaga aatcaaaaga gccatccaag ctaaggacac cttcccaaat 200
gtcactatcc tgtccacatt ggagactctg cagatcatta agcccttaga 250
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 tgccttccca tctaatttat tgtaaagtca tatagtccat gtctgtgatg 850
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<210> 84
<211> 177
<212> PRT
<213> Homo Sapien
<400> 84
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                                                          15
                                      10
   1
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<223> Synthetic oligonucleotide probe

Ile Leu Cys Ser Val Asp Asn His Gly Leu Arg Arg Cys Leu Ile Ser Thr Asp Met His His Ile Glu Glu Ser Phe Gln Glu Ile Lys Arg Ala Ile Gln Ala Lys Asp Thr Phe Pro Asn Val. Thr Ile Leu Ser Thr Leu Glu Thr Leu Gln Ile Ile Lys Pro Leu Asp Val Cys Cys Val Thr Lys Asn Leu Leu Ala Phe Tyr Val Asp Arg Val Phe Lys Asp His Gln Glu Pro Asn Pro Lys Ile Leu Arg Lys Ile Ser Ser Ile Ala Asn Ser Phe Leu Tyr Met Gln Lys Thr Leu Arg Gln Cys Gln Glu Gln Arg Gln Cys His Cys Arg Gln Glu Ala Thr Asn Ala Thr Arg Val Ile His Asp Asn Tyr Asp Gln Leu Glu Val His Ala Ala Ala Ile Lys Ser Leu Gly Glu Leu Asp Val Phe Leu Ala Trp Ile Asn Lys Asn His Glu Val Met Phe Ser Ala

<400> 85

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geeeggggag ggggetgggg etggggeegg aggeggggt tgagtgggtg 200
tgtgeggggg geggaggett gatgcaatee egataagaaa tgetegggtg 250
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agggeeaega ceateeeaae eeggeaetea eageeeegea gegeateeeg 400
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<210> 85

<211> 2137

<212> DNA

<213> Homo Sapien

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cctgtacacc	teeggeeeee	acgggctctc	cagctgcttc	ctgcgcatcc	650
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cagcgtgcgg	tacctctgca	tgggcgccga	cggcaagatg	caggggctgc	800
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cagtgccaaa	cagcggcagc	tgtacaagaa	cagaggcttt	cttccactct	950
ctcatttcct	gcccatgctg	cccatggtcc	cagaggagcc	tgaggacctc	1000
aggggccact	tggaatctga	catgttctct	tcgcccctgg	agaccgacag	1050
catggaccca	tttgggcttg	tcaccggact	ggaggccgtg	aggagtccca	1100
gctttgagaa	gtaactgaga	ccatgcccgg	gcctcttcac	tgctgccagg	1150
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agtccacgtt	ctgtttagct	ttaggaagaa	acatctagaa	gttgtacata	1250
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<210> 86

<211> 216

<212> PRT

<213> Homo Sapien

<400> 86

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Leu Trp Leu Ala Val Ala Gly Arg Pro Leu Ala Phe Ser Asp Ala 20 25 30

Gly Pro His Val His Tyr Gly Trp Gly Asp Pro Ile Arg Leu Arg
35 40 45

His Leu Tyr Thr Ser Gly Pro His Gly Leu Ser Ser Cys Phe Leu 50 55 60

Arg Ile Arg Ala Asp Gly Val Val Asp Cys Ala Arg Gly Gln Ser 65 70 75

Ala His Ser Leu Leu Glu Ile Lys Ala Val Ala Leu Arg Thr Val 80 85 90

Ala Ile Lys Gly Val His Ser Val Arg Tyr Leu Cys Met Gly Ala 95 100 105

Asp Gly Lys Met Gln Gly Leu Leu Gln Tyr Ser Glu Glu Asp Cys
110 115 120

Ala Phe Glu Glu Glu Ile Arg Pro Asp Gly Tyr Asn Val Tyr Arg 125 130 135

Ser Glu Lys His Arg Leu Pro Val Ser Leu Ser Ser Ala Lys Gln
140 145 150

Arg Gln Leu Tyr Lys Asn Arg Gly Phe Leu Pro Leu Ser His Phe
155 160 165

Leu Pro Met Leu Pro Met Val Pro Glu Glu Pro Glu Asp Leu Arg 170 175 180

Gly His Leu Glu Ser Asp Met Phe Ser Ser Pro Leu Glu Thr Asp 185 190 195

Ser Met Asp Pro Phe Gly Leu Val Thr Gly Leu Glu Ala Val Arg 200 205 210

Ser Pro Ser Phe Glu Lys

215

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<223> Synthetic oligonucleotide probe
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<213> Artificial Sequence
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<223> Synthetic oligonucleotide probe
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<210> 89
<211> 22
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<400> 89
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<210> 90
<211> 1857
<212> DNA
<213> Homo Sapien
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 tggcgatcct gttgtgctcc ctggcattgg gcagtgttac agtgcactct 150
 tctgaacctg aagtcagaat tcctgagaat aatcctgtga agttgtcctg 200
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 aagcctacag ttaacatccc ctcctctgcc accattggga accgggcagt 500
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tactggaaat	acaaagttag	ccaggcatgg	tggtgcatgc	ctgtagtccc	1800
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<212> PRT <213> Homo Sapien

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<211> 20
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<223> Synthetic oligonucleotide probe
<400> 94
acacctggtt caaagatggg 20
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<212> DNA
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<223> Synthetic oligonucleotide probe
<400> 95
taggaagagt tgctgaaggc acgg 24
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gtgagggacc agggcgccat gaccgaccag ctgagcaggc ggcagatccg 150
cgagtaccaa ctctacagca ggaccagtgg caagcacgtg caggtcaccg 200
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<210> 99

<211> 205

<212> PRT

<213> Homo Sapien

<400> 99

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20 25 30

Ala Met Thr Asp Gln Leu Ser Arg Arg Gln Ile Arg Glu Tyr Gln
35 40 45

Leu Tyr Ser Arg Thr Ser Gly Lys His Val Gln Val Thr Gly Arg
50 55 60

Arg Ile Ser Ala Thr Ala Glu Asp Gly Asn Lys Phe Ala Lys Leu
65 70 75

Ile Val Glu Thr Asp Thr Phe Gly Ser Arg Val Arg Ile Lys Gly
80 85 90

Ala Glu Ser Glu Lys Tyr Ile Cys Met Asn Lys Arg Gly Lys Leu
95 100 105

Ile Gly Lys Pro Ser Gly Lys Ser Lys Asp Cys Val Phe Thr Glu
110 115 120

Ile Val Leu Glu Asn Asn Tyr Thr Ala Phe Gln Asn Ala Arg His
125 130 135

Glu Gly Trp Phe Met Ala Phe Thr Arg Gln Gly Arg Pro Arg Gln 140 145 150

Ala Ser Arg Ser Arg Gln Asn Gln Arg Glu Ala His Phe Ile Lys 155 160 165

Arg Leu Tyr Gln Gly Gln Leu Pro Phe Pro Asn His Ala Glu Lys
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Gln Lys Gln Phe Glu Phe Val Gly Ser Ala Pro Thr Arg Arg Thr 185 190 195

Lys Arg Thr Arg Arg Pro Gln Pro Leu Thr 200 205

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<210> 101
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 aaaaatgcac aattctatct cttgggcaat cttcacgggg ctggctgctc 200
 tgtgtctctt ccaaggagtg cccgtgcgca gcggagatgc caccttcccc 250
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 gtgcactatt gacaaccggg tcacccgggt ggcctggcta aaccgcagca 350
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ctgcatagca actggtagac cagagcctac ggttacttgg agacacatct 650
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gaaatttgag ggagggaac aaagaatact ttggggggaa aagagtttta 1350
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cttttcccaa acgggaagaa cacagcacac ccggcttgga cccactgcaa 1450
getgeategt geaacetett tggtgecagt gtgggeaagg geteageete 1500
tetgeceaca gagtgecece acgtggaaca ttetggaget ggecatecea 1550
aattcaatca gtccatagag acgaacagaa tgagaccttc cggcccaagc 1600
gtggcgctgc gggcactttg gtagactgtg ccaccacggc gtgtgttgtg 1650
aaacgtgaaa taaaaagagc aaaaaaaaa 1679
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<210> 104

<211> 344

<212> PRT

<213> Homo Sapien

<400>	> 104	1												
Met 1	Lys	Thr	Ile	Gln 5	Pro	Lys	Met	His	Asn 10	Ser	Ile	Ser	Trp	Ala 15
Ile	Phe	Thr	Gly	Leu 20	Ala	Ala	Leu	Cys	Leu 25	Phe	Gln	Gly	Val	Pro 30
Val	Arg	Ser	Gly	Asp 35	Ala	Thr	Phe	Pro	Lys 40	Ala	Met	Asp	Asn	Val 45
Thr	Val	Arg	Gln	Gly 50	Glu	Ser	Ala	Thr	Leu 55	Arg	Cys	Thr	Ile	Asp 60
Asn	Arg	Val	Thr	Arg 65	Val	Ala	Trp	Leu	Asn 70	Arg	Ser	Thr	Ile	Leu 75
Tyr	Ala	Gly	Asn	Asp 80	Lys	Trp	Cys	Leu	Asp 85	Pro	Arg	Val	Val	Leu 90
Leu	Ser	Asn	Thr	Gln 95	Thr	Gln	Tyr	Ser	Ile 100	Glu	Ile	Gln	Asn	Val 105
Asp	Val	Tyr	Asp	Glu 110	Gly	Pro	Tyr	Thr	Cys 115	Ser	Val	Gln	Thr	Asp 120
Asn	His	Pro	Lys	Thr 125	Ser	Arg	Val	His	Leu 130	Ile	Val	Gln	Val	Ser 135
Pro	Lys	Ile	Val	Glu 140		Ser	Ser	Asp	Ile 145	Ser	Ile	Asn	Glu	Gly 150
Asn	Asn	Ile	Ser	Leu 155	Thr	Cys	Ile	Ala	Thr 160	Gly	Arg	Pro	Glu	Pro 165
Thr	Val	Thr	Trp	Arg 170	His	Ile	Ser	Pro	Lys 175	Ala	Val	Gly	Phe	Val 180
Ser	Glu	Asp	Glu	Tyr 185	Leu	Glu	Ile	Gln	Gly 190	Ile	Thr	Arg	Glu	Gln 195
Ser	Gly	Asp	Tyr	Glu 200	Cys	Ser	Ala	Ser	Asn 205	Asp	Val	Ala	Ala	Pro 210
Val	Val	Arg	Arg	Val 215	Lys	Val	Thr	Val	Asn 220	Tyr	Pro	Pro	Tyr	Ile 225
Ser	Glu	Ala	Lys	Gly 230	Thr	Gly	Val	Pro	Val 235	Gly	Gln	Lys	Gly	Thr 240
Leu	Gln	Cys	Glu	Ala 245	Ser	Ala	Val	Pro	Ser 250	Ala	Glu	Phe	Gln	Trp 255
Tyr	Lys	Asp	Asp	Lys 260	Arg	Leu	Ile	Glu	Gly 265	Lys	Lys	Gly	Val	Lys 270
Val	Glu	Asn	Aṛg	Pro 275	Phe	Leu	Ser	_Lys_	Leu 280	Ile	Phe	Phe	Asn	Val 285

Ser Glu His Asp Tyr Gly Asn Tyr Thr Cys Val Ala Ser Asn Lys 290 295 300

Leu Gly His Thr Asn Ala Ser Ile Met Leu Phe Gly Pro Gly Ala 305 310

Val Ser Glu Val Ser Asn Gly Thr Ser Arg Arg Ala Gly Cys Val 320 325 330

Trp Leu Leu Pro Leu Leu Val Leu His Leu Leu Leu Lys Phe 335

- <210> 105
- <211> 1734
- <212> DNA
- <213> Homo Sapien

<400> 105

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<210> 106

<211> 440

<212> PRT

<213> Homo Sapien

<400> 106

Met Lys Phe Gln Gly Pro Leu Ala Cys Leu Leu Leu Ala Leu Cys
1 10 15

Leu Gly Ser Gly Glu Ala Gly Pro Leu Gln Ser Gly Glu Glu Ser

Thr Gly Thr Asn Ile Gly Glu Ala Leu Gly His Gly Leu Gly Asp

Ala Leu Ser Glu Gly Val Gly Lys Ala Ile Gly Lys Glu Ala Gly
50 55 60

Gly Ala Ala Gly Ser Lys Val Ser Glu Ala Leu Gly Gln Gly Thr
65 70 75

Arg Glu Ala Val Gly Thr Gly Val Arg Gln Val Pro Gly Phe Gly 80 85 90

Leu Gly Asn Thr Gly His Glu Ile Gly Arg Gln Ala Glu Asp Val

				110					115					120
Ile	Arg	His	Gly	Ala 125	Asp	Ala	Val	Arg	Gly 130	Ser	Trp	Gln	Gly	Val 135
Pro	Gly	His	Ser	Gly 140	Ala	Trp	Glu	Thr	Ser 145	Gly	Gly	His	Gly	Ile 150
Phe	Gly	Ser	Gln	Gly 155	Gly	Leu	Gly	Gly	Gln 160	Gly	Gln	Gly	Asn	Pro 165
Gly	Gly	Leu	Gly	Thr 170	Pro	Trp	Val	His	Gly 175	Tyr	Pro	Gly	Asn	Ser 180
Ala	Gly	Ser	Phe	Gly 185	Met	Asn	Pro	Gln	Gly 190	Ala	Pro	Trp	Gly	Gln 195
Gly	Gly	Asn	Gly	Gly 200	Pro	Pro	Asn	Phe	Gly 205	Thr	Asn	Thr	Gln	Gly 210
Ala	Val	Ala	Gln	Pro 215	Gly	Tyr	Gly	Ser	Val 220	Arg	Ala	Ser	Asn	Gln 225
Asn	Glu	Gly	Cys	Thr 230	Asn	Pro	Pro	Pro	Ser 235	Gly	Ser	Gly	Gly	Gly 240
Ser	Ser	Asn	Ser	Gly 245	Gly	Gly	Ser	Gly	Ser 250	Gln	Ser	Gly	Ser	Ser 255
Gly	Ser	Gly	Ser	Asn 260	Gly	Asp	Asn	Asn	Asn 265	Gly	Ser	Ser	Ser	Gly 270
Gly	Ser	Ser	Ser	Gly 275	Ser	Ser	Ser	Gly	Ser 280	Ser	Ser	Gly	Gly	Ser 285
Ser	Gly	Gly	Ser	Ser 290	Gly	Gly	Ser	Ser	Gly 295	Asn	Ser	Gly	Gly	Ser 300
Arg	Gly	Asp	Ser	Gly 305	Ser	Glu	Ser	Ser	Trp 310	Gly	Ser	Ser	Thr	Gly 315
Ser	Ser	Ser	Gly	Asn 320	His	Gly	Gly	Ser	Gly 325	Gly	Gly	Asn	Gly	His 330
Lys	Pro	Gly	Cys	Glu 335	Lys	Pro	Gly	Asn	Glu 340	Ala	Arg	Gly	Ser	Gly 345
Glu	Ser	Gly	Ile	Gln 350	Gly	Phe	Arg	Gly	Gln 355	Gly	Val	Ser	Ser	Asn 360
Met	Arg	Glu	Ile	Ser 365	Lys	Glu	Gly	Asn	Arg 370	Leu	Leu	Gly	Gly	Ser 375
Gly	Asp	Asn	Tyr	Arg 380	Gly	Gln	Gly	Ser	Ser 385	Trp	Gly	Ser	Gly	Gly 390
Gly	Asp	Ala	Val	Gly 395	Gly	Val	Asn	Thr	Val 400	Asn	Ser	Glu	Thr	Ser 405

Pro Gly Met Phe Asn Phe Asp Thr Phe Trp Lys Asn Phe Lys Ser 410 415 420

Lys Leu Gly Phe Ile Asn Trp Asp Ala Ile Asn Lys Asp Gln Arg 425 430 435

Ser Ser Arg Ile Pro 440

<210> 107

<211> 918

<212> DNA

<213> Homo Sapien

<400> 107 agccaggcag cacatcacag cgggaggagc tgtcccaggt ggcccagctc 50 ctgcgctctg cctgacaggg tcccaagccc tgcagtgcta cagctttgag 150 cacacctact ttggcccctt tgacctcagg gccatgaagc tgcccagcat 200 ctcctgtcct catgagtgct ttgaggctat cctgtctctg gacaccgggt 250 ategegegee ggtgaeeetg gtgeggaagg getgetggae egggeeteet 300 gcgggccaga cgcaatcgaa cccggacgcg ctgccgccag actactcggt 350 ggtgcgcgc tgcacaactg acaaatgcaa cgcccacctc atgactcatg 400 acgccctccc caacctgagc caagcacccg acccgccgac gctcagcggc 450 geogagtget aegeetgtat eggggteeae eaggatgaet gegetategg 500 caggtcccga cgagtccagt gtcaccagga ccagaccgcc tgcttccagg 550 gcagtggcag aatgacagtt ggcaatttct cagtccctgt gtacatcaga 600 acctgccacc ggccctcctg caccaccgag ggcaccacca gcccctggac 650 agccatcgac ctccagggct cctgctgtga ggggtacctc tgcaacagga 700 aatccatgac ccagcccttc accagtgctt cagccaccac ccctccccga 750 gcactacagg teetggeeet geteeteeca gteeteetge tggtgggget 800 ctcagcatag accgccctc caggatgctg gggacagggc tcacacacct 850 cattettqct getteagece etateacata geteactgga aaatgatgtt 900 aaagtaagaa ttgcaaaa 918

<210> 108

<211> 251

<212> PRT

<213> Homo Sapien

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<400> 108
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Ala Ala Leu Cys Leu Thr Gly Ser Gln Ala Leu Gln Cys Tyr Ser
                                                            30
                  20
                                       25
Phe Glu His Thr Tyr Phe Gly Pro Phe Asp Leu Arg Ala Met Lys
                                                            45
Leu Pro Ser Ile Ser Cys Pro His Glu Cys Phe Glu Ala Ile Leu
                  50
                                       55
 Ser Leu Asp Thr Gly Tyr Arg Ala Pro Val Thr Leu Val Arg Lys
                  65
                                       70
                                                            75
Gly Cys Trp Thr Gly Pro Pro Ala Gly Gln Thr Gln Ser Asn Pro
                                                            90
                                       85
Asp Ala Leu Pro Pro Asp Tyr Ser Val Val Arg Gly Cys Thr Thr
                                                           105
                  95
Asp Lys Cys Asn Ala His Leu Met Thr His Asp Ala Leu Pro Asn
                 110
                                      115
                                                           120
Leu Ser Gln Ala Pro Asp Pro Pro Thr Leu Ser Gly Ala Glu Cys
                 125
                                                          135
                                      130
Tyr Ala Cys Ile Gly Val His Gln Asp Asp Cys Ala Ile Gly Arg
                                                           150
                 140
                                      145
Ser Arg Arg Val Gln Cys His Gln Asp Gln Thr Ala Cys Phe Gln
                 155
                                      160
                                                           165
Gly Ser Gly Arg Met Thr Val Gly Asn Phe Ser Val Pro Val Tyr
                 170
                                      175
                                                           180
Ile Arg Thr Cys His Arg Pro Ser Cys Thr Thr Glu Gly Thr Thr
                                                          195
                                      190
                 185
             Thr Ala Ile Asp Leu Gln Gly Ser Cys Cys Glu Gly
                 200
                                      205
Tyr Leu Cys Asn Arg Lys Ser Met Thr Gln Pro Phe Thr Ser Ala
                                      220
                 215
Ser Ala Thr Thr Pro Pro Arg Ala Leu Gln Val Leu Ala Leu Leu
                                                           240
                                      235
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Leu Pro Val Leu Leu Val Gly Leu Ser Ala
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                                      250
<210> 109
<211> 1813
<212> DNA
<213> Homo Sapien
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<400> 109

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taagacctaa gtgtccagga gacagaagga gaagaggaag tggatctgga 1500 attgggagga gcctccaccc acccetgact cetecttatg aagccagctg 1550 ctgaaattag ctactcacca agagtgaggg gcagagactt ccagtcactg 1600 agtctccag gccccttga tetgtacccc acccetatet aacaccaccc 1650 ttggctccca ctccagctcc ctgtattgat ataacctgtc aggctggctt 1700 ggttaggttt tactggggca gaggataggg aatctcttat taaaactaac 1750 atgaaatatg tgttgtttc atttgcaaat ttaaataaag atacataatg 1800 tttgtatgaa aaa 1813

<210> 110

<211> 390

<212> PRT

<213> Homo Sapien

<400> 110

Met Ile Ser Leu Pro Gly Pro Leu Val Thr Asn Leu Leu Arg Phe
1 5 10 15

Leu Phe Leu Gly Leu Ser Ala Leu Ala Pro Pro Ser Arg Ala Gln
20 25 30

Leu Gln Leu His Leu Pro Ala Asn Arg Leu Gln Ala Val Glu Gly
35 40 45

Gly Glu Val Val Leu Pro Ala Trp Tyr Thr Leu His Gly Glu Val
50 55 60

Ser Ser Ser Gln Pro Trp Glu Val Pro Phe Val Met Trp Phe Phe 65 70 75

Lys Gln Lys Glu Lys Glu Asp Gln Val Leu Ser Tyr Ile Asn Gly 80 85 90

Val Thr Thr Ser Lys Pro Gly Val Ser Leu Val Tyr Ser Met Pro 95 100 105

Ser Arg Asn Leu Ser Leu Arg Leu Glu Gly Leu Gln Glu Lys Asp 110 115 120

Ser Gly Pro Tyr Ser Cys Ser Val Asn Val Gln Asp Lys Gln Gly 125 130 135

Lys Ser Arg Gly His Ser Ile Lys Thr Leu Glu Leu Asn Val Leu 140 145 150

Val Pro Pro Ala Pro Pro Ser Cys Arg Leu Gln Gly Val Pro His
155 160 165

Val Gly Ala Asn Val Thr Leu Ser Cys Gln Ser Pro Arg Ser Lys
170 175 180

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Pro Ala Val Gln Tyr Gln Trp Asp Arg Gln Leu Pro Ser Phe Gln
                                      190
                                                           195
                  185
 Thr Phe Phe Ala Pro Ala Leu Asp Val Ile Arg Gly Ser Leu Ser
                                       205
                                                           210
                  200
 Leu Thr Asn Leu Ser Ser Ser Met Ala Gly Val Tyr Val Cys Lys
                  215
                                      220
                                                           225
 Ala His Asn Glu Val Gly Thr Ala Gln Cys Asn Val Thr Leu Glu
                  230
                                      235
                                                           240
 Val Ser Thr Gly Pro Gly Ala Ala Val Val Ala Gly Ala Val Val
                                                           255
                  245
                                       250
 Gly Thr Leu Val Gly Leu Gly Leu Leu Ala Gly Leu Val Leu Leu
                                      265
                                                           270
                  260
 Tyr His Arg Arg Gly Lys Ala Leu Glu Glu Pro Ala Asn Asp Ile
                  275
                                      280
                                                           285
 Lys Glu Asp Ala Ile Ala Pro Arg Thr Leu Pro Trp Pro Lys Ser
                                                           300
                  290
                                      295
 Ser Asp Thr Ile Ser Lys Asn Gly Thr Leu Ser Ser Val Thr Ser
                  305
                                      310
                                                           315
 Ala Arg Ala Leu Arg Pro Pro His Gly Pro Pro Arg Pro Gly Ala
                  320
                                      325
                                                           330
 Leu Thr Pro Thr Pro Ser Leu Ser Ser Gln Ala Leu Pro Ser Pro
                  335
                                       340
                                                           345
 Arg Leu Pro Thr Thr Asp Gly Ala His Pro Gln Pro Ile Ser Pro
                  350
                                      355
                                                           360
 Ile Pro Gly Gly Val Ser Ser Ser Gly Leu Ser Arg Met Gly Ala
                  365
                                      370
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 Val Pro Val Met Val Pro Ala Gln Ser Gln Ala Gly Ser Leu Val
<210> 111
<211> 22
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 111
 agggteteca ggagaaagae te 22
<210> 112
<211> 24
_<212> DNA
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<213> Artificial Sequence

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<220>
<223> Synthetic oligonucleotide probe
<400> 112
attgtgggcc ttgcagacat agac 24
<210> 113
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic oligonucleotide probe
<400> 113
ggccacagca tcaaaacctt agaactcaat gtactggttc ctccagctcc 50
<210> 114
<211> 2479
<212> DNA
<213> Homo Sapien
<400> 114
acttgccatc acctgttgcc agtgtggaaa aattctccct gttgaatttt 50
ttgcacatgg aggacagcag caaagagggc aacacaggct gataagacca 100
gagacagcag ggagattatt ttaccatacg ccctcaggac gttccctcta 150
getggagtte tggaetteaa cagaaceeea teeagteatt ttgattttge 200
ccgtacttca gaaatgggcc tacagaccac aaagtggccc agccatgggg 300
cttttttcct gaagtcttgg cttatcattt ccctggggct ctactcacag 350
 gtgtccaaac tcctggcctg ccctagtgtg tgccgctgcg acaggaactt 400
 tgtctactgt aatgagcgaa gcttgacctc agtgcctctt gggatcccgg 450
 agggcgtaac cgtactctac ctccacaaca accaaattaa taatgctgga 500
 tttcctgcag aactgcacaa tgtacagtcg gtgcacacgg tctacctgta 550
 tggcaaccaa ctggacgaat tccccatgaa ccttcccaag aatgtcagag 600
 ttctccattt gcaggaaaac aatattcaga ccatttcacg ggctgctctt 650
geceagetet tgaagettga agagetgeae etggatgaea aetecatate 700
 cacagtgggg gtggaagacg gggccttccg ggaggctatt agcctcaaat 750
 tgttgttttt gtctaagaat cacctgagca gtgtgcctgt tgggcttcct 800
gtggacttgc aagagctgag agtggatgaa aatcgaattg ctgtcatatc 850
 cgacatggcc ttccagaatc tcacgagctt ggagcgtctt attgtggacg 900
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ggaacctcct gaccaacaag ggtatcgccg agggcacctt cagccatctc 950 accaagetea aggaatttte aattgtaegt aattegetgt eccaeeetee 1000 tecegatete ceaggtacge atetgateag getetatttg caggacaace 1050 agataaacca cattcctttg acagccttct caaatctgcg taagctggaa 1100 cggctggata tatccaacaa ccaactgcgg atgctgactc aaggggtttt 1150 tgataatete tecaacetga ageageteae tgeteggaat aaceettggt 1200 tttgtgactg cagtattaaa tgggtcacag aatggctcaa atatatccct 1250 tcatctctca acgtgcgggg tttcatgtgc caaggtcctg aacaagtccg 1300 ggggatggcc gtcagggaat taaatatgaa tcttttgtcc tgtcccacca 1350 cgaccccgg cctgcctctc ttcaccccag ccccaagtac agcttctccg 1400 accactcage eteceacet etetatteca aaccetagea gaagetacae 1450 gcctccaact cctaccacat cgaaacttcc cacgattcct gactgggatg 1500 gcagagaaag agtgacccca cctatttctg aacggatcca gctctctatc 1550 cattttgtga atgatacttc cattcaagtc agctggctct ctctcttcac 1600 cgtgatggca tacaaactca catgggtgaa aatgggccac agtttagtag 1650 ggggcatcgt tcaggagcgc atagtcagcg gtgagaagca acacctgagc 1700 ctggttaact tagagccccg atccacctat cggatttgtt tagtgccact 1750 ggatgctttt aactaccgcg cggtagaaga caccatttgt tcagaggcca 1800 ccacccatgc ctcctatctg aacaacggca gcaacacagc gtccagccat 1850 gagcagacga cgtcccacag catgggctcc ccctttctgc tggcgggctt 1900 gateggggge geggtgatat ttgtgetggt ggtettgete agegtetttt 1950 gctggcatat gcacaaaaag gggcgctaca cctcccagaa gtggaaatac 2000 aaccggggcc ggcggaaaga tgattattgc gaggcaggca ccaagaagga 2050 caactccatc ctggagatga cagaaaccag ttttcagatc gtctccttaa 2100 ataacgatca actccttaaa ggagatttca gactgcagcc catttacacc 2150 ccaaatgggg gcattaatta cacagactgc catatcccca acaacatgcg 2200 atactgcaac agcagcgtgc cagacctgga gcactgccat acgtgacagc 2250 cagaggccca gcgttatcaa ggcggacaat tagactcttg agaacacact 2300 cgtgtgtgca cataaagaca cgcagattac atttgataaa tgttacacag 2350 atgcatttgt gcatttgaat actctgtaat ttatacggtg tactatata 2400 tgggatttaa aaaaagtgct atcttttcta tttcaagtta attacaaaca 2450 gttttgtaac tctttgcttt ttaaatctt 2479

- <210> 115
- <211> 660
- <212> PRT
- <213> Homo Sapien
- <400> 115
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- Leu Lys Ser Trp Leu Ile Ile Ser Leu Gly Leu Tyr Ser Gln Val 20 25 30
- Ser Lys Leu Leu Ala Cys Pro Ser Val Cys Arg Cys Asp Arg Asn 35 40 45
- Phe Val Tyr Cys Asn Glu Arg Ser Leu Thr Ser Val Pro Leu Gly 50 55 60
- Ile Pro Glu Gly Val Thr Val Leu Tyr Leu His Asn Asn Gln Ile
 65 70 75
- Asn Asn Ala Gly Phe Pro Ala Glu Leu His Asn Val Gln Ser Val
 80 85 90
- His Thr Val Tyr Leu Tyr Gly Asn Gln Leu Asp Glu Phe Pro Met
 95 100 105
- Asn Leu Pro Lys Asn Val Arg Val Leu His Leu Gln Glu Asn Asn 110 115 120
- Ile Gln Thr Ile Ser Arg Ala Ala Leu Ala Gln Leu Leu Lys Leu 125 130 135
- Glu Glu Leu His Leu Asp Asp Asn Ser Ile Ser Thr Val Gly Val
 140 145 150
- Glu Asp Gly Ala Phe Arg Glu Ala Ile Ser Leu Lys Leu Leu Phe 155 160 165
- Leu Ser Lys Asn His Leu Ser Ser Val Pro Val Gly Leu Pro Val
 170 175 180
- Asp Leu Gln Glu Leu Arg Val Asp Glu Asn Arg Ile Ala Val Ile 185 190 195
- Ser Asp Met Ala Phe Gln Asn Leu Thr Ser Leu Glu Arg Leu Ile 200 205 210
- Val Asp Gly Asn Leu Leu Thr Asn Lys Gly Ile Ala Glu Gly Thr
- Phe Ser His Leu Thr Lys Leu Lys Glu Phe Ser Ile Val Arg Asn

				230					235					240
Ser	Leu	Ser	His	Pro 245	Pro	Pro	Asp	Leu	Pro 250	Gly	Thr	His	Leu	Ile 255
Arg	Leu	Tyr	Leu	Gln 260	Asp	Asn	Gln	Ile	Asn 265	His	Ile	Pro	Leu	Thr 270
Ala	Phe	Ser	Asn	Leu 275	Arg	Lys	Leu	Glu	Arg 280	Leu	Asp	Ile	Ser	Asn 285
Asn	Gln	Leu	Arg	Met 290	Leu	Thr	Gln	Gly	Val 295	Phe	Asp	Asn	Leu	Ser 300
Asn	Leu	Lys	Gln	Leu 305	Thr	Ala	Arg	Asn	Asn 310	Pro	Trp	Phe	Cys	Asp 315
Cys	Ser	Ile	Lys	Trp 320	Val	Thr	Glu	Trp	Leu 325	Lys	Tyr	Ile	Pro	Ser 330
Ser	Leu	Asn	Val	Arg 335	Gly	Phe	Met	Cys	Gln 340	Gly	Pro	Glu	Gln	Val 345
Arg	Gly	Met	Ala	Val 350	Arg	Glu	Leu	Asn	Met 355	Asn	Leu	Leu	Ser	Cys 360
Pro	Thr	Thr	Thr	Pro 365	Gly	Leu	Pro	Leu	Phe 370	Thr	Pro	Ala	Pro	Ser 375
Thr	Ala	Ser	Pro	Thr 380	Thr	Gln	Pro	Pro	Thr 385	Leu	Ser	Ile	Pro	Asn 390
Pro	Ser	Arg	Ser	Tyr 395	Thr	Pro	Pro	Thr	Pro 400	Thr	Thr	Ser	Lys	Leu 405
Pro	Thr	Ile	Pro	Asp 410	Trp	Asp	Gly	Arg	Glu 415	Arg	Val	Thr	Pro	Pro 420
Ile	Ser	Glu	Arg	Ile 425	Gln	Leu	Ser	Ile	His 430	Phe	Val	Asn	Asp	Thr 435
Ser	Ile	Gln	Val	Ser 440	Trp	Leu	Ser	Leu	Phe 445	Thr	Val	Met	Ala	Tyr 450
Lys	Leu	Thr	Trp	Val 455	Lys	Met	Gly	His	Ser 460	Leu	Val	Gly	Gly	Ile 465
Val	Gln	Glu	Arg	Ile 470	Val	Ser	Gly	Glu	Lys 475	Gln	His	Leu	Ser	Leu 480
Val	Asn	Leu	Glu	Pro 485	Arg	Ser	Thr	Tyr	Arg 490	Ile	Cys	Leu	Val	Pro 495
Leu	Asp	Ala	Phe	Asn 500	Tyr	Arg	Ala	Val	Glu 505	Asp	Thr	Ile	Cys	Ser 510
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                                                           555
 Val Val Leu Leu Ser Val Phe Cys Trp His Met His Lys Lys Gly
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 Asp Asp Tyr Cys Glu Ala Gly Thr Lys Lys Asp Asn Ser Ile Leu
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 Glu Met Thr Glu Thr Ser Phe Gln Ile Val Ser Leu Asn Asp
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<212> PRT

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Val Trp Asn Gln Phe Phe Val Pro Glu Glu Met Asn Thr Thr Ser 50 55 60

His His Ile Gly Gln Leu Arg Ser Asp Leu Asp Asn Gly Asn Asn 65 70 75

Ser Phe Gln Tyr Lys Leu Leu Gly Ala Gly Ala Gly Ser Thr Phe 80 85 90

Ile Ile Asp Glu Arg Thr Gly Asp Ile Tyr Ala Ile Gln Lys Leu 95 100 105

Asp Arg Glu Glu Arg Ser Leu Tyr Ile Leu Arg Ala Gln Val Ile 110 115 120

Asp Ile Ala Thr Gly Arg Ala Val Glu Pro Glu Ser Glu Phe Val 125 130 135

Ile Lys Val Ser Asp Ile Asn Asp Asn Glu Pro Lys Phe Leu Asp 140 145 150

Glu Pro Tyr Glu Ala Ile Val Pro Glu Met Ser Pro Glu Gly Thr

Leu Val Ile Gln Val Thr Ala Ser Asp Ala Asp Asp Pro Ser Ser

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Tyr	Phe	Ser	Val	Glu 200	Pro	Thr	Thr	Gly	Val 205	Ile	Arg	Ile	Ser	Ser 210
Lys	Met	Asp	Arg	Glu 215	Leu	Gln	Asp	Glu	Tyr 220	Trp	Val	Ile	Ile	Gln 225
Ala	Lys	Asp	Met	Ile 230	Gly	Gln	Pro	Gly	Ala 235	Leu	Ser	Gly	Thr	Thr 240
Ser	Val	Leu	Ile	Lys 245	Leu	Ser	Asp	Val	Asn 250	Asp	Asn	Lys	Pro	Ile 255
Phe	Lys	Glu	Ser	Leu 260	Tyr	Arg	Leu	Thr	Val 265	Ser	Glu	Ser	Ala	Pro 270
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Gly	Glu	Asn	Ala	Glu 290	Met	Asp	Tyr	Ser	Ile 295	Glu	Glu	Asp	Asp	Ser 300
Gln	Thr	Phe	Asp	Ile 305	Ile	Thr	Asn	His	Glu 310	Thr	Gln	Glu	Gly	Ile 315
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Gly	Ile	Arg	Ala	Lys 335	Val	Lys	Asn	His	His 340	Val	Pro	Glu	Gln	Leu 345
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Val	Val	Ser	Ala	Thr 395	Asp	Pro	Asp	Asn	Arg 400	Lys	Ser	Pro	Ile	Arg 405
Tyr	Ser	Ile	Thr	Arg 410	Ser	Lys	Val	Phe	Asn 415	Ile	Asn	Asp	Asn	Gly 420
Thr	Ile	Thr	Thr	Ser 425	Asn	Ser	Leu	Asp	Arg 430	Glu	Ile	Ser	Ala	Trp 435
Tyr	Asn	Leu	Ser	Ile 440	Thr	Ala	Thr	Glu	Lys 445	Tyr	Asn	Ile	Glu	Gln 450
Ile	Ser	Ser	Ile	Pro 455	Leu	Tyr	Val	Gln	Val 460	Leu	Asn	Ile	Asn	Asp 465

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Gln	Glu	Glu	Pro	Val 545	Phe	Tyr	Ile	Ser	Ile 550	Leu	Ile	Ala	Asp	Asn 555
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Cys	Asp	Cys	Gly	Asp 575	Ser	Gly	Ser	Thr	Gln 580	Thr	Cys	Gln	Tyr	Gln 585
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Ile	Leu	Ile	Cys	Ile 605	Met	Ile	Ile	Phe	Gly 610	Phe	Ile	Phe	Leu	Thr 615
Leu	Gly	Leu	Lys	Gln 620	Arg	Arg	Lys	Gln	Ile 625	Leu	Phe	Pro	Glu	Lys 630
Ser	Glu	Asp	Phe	Arg 635	Glu	Asn	Ile	Phe	Gln 640	Tyr	Asp	Asp	Glu	Gly 645
Gly	Gly	Glu	Glu	Asp 650	Thr	Glu	Ala	Phe	Asp 655	Ile	Ala	Glu	Leu	Arg 660
Ser	Ser	Thr	Ile	Met 665	Arg	Glu	Arg	Lys	Thr 670	Arg	Lys	Thr	Thr	Ser 675
Ala	Glu	Ile	Arg	Ser 680	Leu	Tyr	Arg	Gln	Ser 685	Leu	Gln	Val	Gly	Pro 690
Asp	Ser	Ala	Ile	Phe 695	Arg	Lys	Phe	Ile	Leu 700	Glu	Lys	Leu	Glu	Glu 705
Ala	Asn	Thr	Asp	Pro 710	Cys	Ala	Pro	Pro	Phe 715	Asp	Ser	Leu	Gln	Thr 720
Tyr	Ala	Phe	Glu	Gly 725	Thr	Gly	Ser	Leu	Ala 730	Gly	Ser	Leu	Ser	Ser 735
Leu			Ala					_			_	_	_	
Asn	Glu	Len	Glv	Pro	Ara	Phe	Lvc	Δra	T.e.i	Δla	Cve	Met	Dhe	G] v

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<211> 179

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<213> Homo Sapien

<400> 126

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Asn Phe Gln Gln Pro Tyr Ile Thr Asn Arg Thr Phe Met Leu Ala 50 55 60

Lys Glu Ala Ser Leu Ala Asp Asn Asn Thr Asp Val Arg Leu Ile
65 70 75

Gly Glu Lys Leu Phe His Gly Val Ser Met Ser Glu Arg Cys Tyr

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<212> DNA

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<212> PRT

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120 110 Tyr Gly Pro Ser Asp Ser Ala Ser Arg Asp Leu Trp Val Asn Ile 130 125 Asp Gln Met Glu Lys Asp Lys Val Lys Ile His Gly Ile Leu Ser 145 150 140 Asn Thr His Arg Gln Ala Ala Arg Val Asn Leu Ser Phe Asp Phe 165 155 160 Pro Phe Tyr Gly His Phe Leu Arg Glu Ile Thr Val Ala Thr Gly

Gly Phe Ile Tyr Thr Gly Glu Val Val His Arg Met Leu Thr Ala 195 190 185

170

175

180

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Val	Val	Gln	Trp	Asp 230	His	Val	His	Leu	Gln 235	Asp	Asn	Tyr	Asn	Leu 240
Gly	Ser	Phe	Thr	Phe 245	Gln	Ala	Thr	Leu	Leu 250	Met	Asp	Gly	Arg	Il∈ 255
Ile	Phe	Gly	Tyr	Lys 260	Glu	Ile	Pro	Val	Leu 265	Val	Thr	Gln	Ile	Ser 270
Ser	Thr	Asn	His	Pro 275	Val	Lys	Val	Gly	Leu 280	Ser	Asp	Ala	Phe	Val 285
Val	Val	His	Arg	Ile 290	Gln	Gln	Ile	Pro	Asn 295	Val	Arg	Arg	Arg	Thr 300
Ile	Tyr	Glu	Tyr	His 305	Arg	Val	Glu	Leu	Gln 310	Met	Ser	Lys	Ile	Thr 315
Asn	Ile	Ser	Ala	Val 320	Glu	Met	Thr	Pro	Leu 325	Pro	Thr	Cys	Leu	Gln 330
Phe	Asn	Arg	Cys	Gly 335	Pro	Cys	Val	Ser	Ser 340	Gln	Ile	Gly	Phe	Asn 345
Cys	Ser	Trp	Cys	Ser 350	Lys	Leu	Gln	Arg	Cys 355	Ser	Ser	Gly	Phe	Asp 360
Arg	His	Arg	Gln	Asp 365	Trp	Val	Asp	Ser	Gly 370	Cys	Pro	Glu	Glu	Ser 375
Lys	Glu	Lys	Met	Cys 380	Glu	Asn	Thr	Glu	Pro 385	Val	Glu	Thr	Ser	Ser 390
Arg	Thr	Thr	Thr	Thr 395	Val	Gly	Ala	Thr	Thr 400	Thr	Gln	Phe	Arg	Val 405
Leu	Thr	Thr	Thr	Arg 410	Arg	Ala	Val	Thr	Ser 415	Gln	Phe	Pro	Thr	Ser 420
Leu	Pro	Thr	Glu	Asp 425	Asp	Thr	Lys	Ile	Ala 430	Leu	His	Leu	Lys	Asp 435
Asn	Gly	Ala	Ser	Thr 440	Asp	Asp	Ser	Ala	Ala 445	Glu	Lys	Lys	Gly	Gly 450
Thr	Leu	His	Ala	Gly 455	Leu	Ile	Ile	Gly	Ile 460	Leu	Ile	Leu	Val	Leu 465
Ile	Val	Ala		Ala 470						_	Met	-		His 480
Dro	Thr	Ser	Δla	Δla	Ser	Tlo	Dho'	Dho	Tla	G 333	7 200	7) 22 0	Dro	802

485 490 495

Arg Trp Pro Ala Met Lys Phe Arg Arg Gly Ser Gly His Pro Ala 500 505 510

Tyr Ala Glu Val Glu Pro Val Gly Glu Lys Glu Gly Phe Ile Val 515 520 525

Ser Glu Gln Cys

<210> 129

<211> 4834

<212> DNA

<213> Homo Sapien

<220>

<221> unsure

<222> 3784

<223> unknown base

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<210> 130

<211> 354

<212> PRT

<213> Homo Sapien

<400> 130

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Trp Leu Ala Ala Val Leu Leu Ser Leu Cys Cys Leu Leu Pro Ser 20 25 30

Cys Leu Pro Ala Gly Gln Ser Val Asp Phe Pro Trp Ala Ala Val

Asp Asn Met Met Val Arg Lys Gly Asp Thr Ala Val Leu Arg Cys

				50					55					60
Tyr	Leu	Glu	Asp	Gly 65	Ala	Ser	Lys	Gly	Ala 70	Trp	Leu	Asn	Arg	Ser 75
Ser	Ile	Ile	Phe	Ala 80	Gly	Gly	Asp	Lys	Trp 85	Ser	Val	Asp	Pro	Arg 90
Val	Ser	Ile	Ser	Thr 95	Leu	Asn	Lys	Arg	Asp 100	Tyr	Ser	Leu	Gln	Ile 105
Gln	Asn	Val	Asp	Val 110	Thr	Asp	Asp	Gly	Pro 115	Tyr	Thr	Cys	Ser	Val 120
Gln	Thr	Gln	His	Thr 125	Pro	Arg	Thr	Met	Gln 130	Val	His	Leu	Thr	Val 135
Gln	Val	Pro	Pro	Lys 140	Ile	Tyr	Asp	Ile	Ser 145	Asn	Asp	Met	Thr	Val 150
Asn	Glu	Gly	Thr	Asn 155	Val	Thr	Leu	Thr	Cys 160	Leu	Ala	Thr	Gly	Lys 165
Pro	Glu	Pro	Ser	Ile 170	Ser	Trp	Arg	His	Ile 175	Ser	Pro	Ser	Ala	Lys 180
Pro	Phe	Glu	Asn	Gly 185	Gln	Tyr	Leu	Asp	Ile 190	Tyr	Gly	Ile	Thr	Arg 195
Asp	Gln	Ala	Gly	Glu 200	Tyr	Glu	Cys	Ser	Ala 205	Glu	Asn	Asp	Val	Ser 210
Phe	Pro	Asp	Val	Arg 215	Lys	Val	Lys	Val	Val 220	Val	Asn	Phe	Ala	Pro 225
Thr	Ile	Gln	Glu	Ile 230	Lys	Ser	Gly	Thr	Val 235	Thr	Pro	Gly	Arg	Ser 240
Gly	Leu	Ile	Arg	Cys 245	Glu	Gly	Ala	Gly	Val 250	Pro	Pro	Pro	Ala	Phe 255
Glu	Trp	Tyr	Lys	Gly 260	Glu	Lys	Lys	Leu	Phe 265	Asn	Gly	Gln	Gln	Gly 270
Ile	Ile	Ile	Gln	Asn 275	Phe	Ser	Thr	Arg	Ser 280	Ile	Leu	Thr	Val	Thr 285
Asn	Val	Thr	Gln	Glu 290	His	Phe	Gly	Asn	Tyr 295	Thr	Cys	Val	Ala	Ala 300
Asn	Lys	Leu	Gly	Thr 305	Thr	Asn	Ala	Ser	Leu 310	Pro	Leu	Asn	Pro	Pro 315
Ser	Thr	Ala	Gln	Tyr 320	Gly	Ile	Thr	Gly	Ser 325	Ala	Asp	Val	Leu	Phe 330
Ser	Cys	Trp	Tyr	Leu 335	Val	Leu	Thr	Leu	Ser 340	Ser	Phe	Thr	Ser	Ile 345

Phe Tyr Leu Lys Asn Ala Ile Leu Gln 350

- <210> 131
- <211> 823
- <212> DNA
- <213> Homo Sapien
- <400> 131

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- <210> 132
- <211> 155
- <212> PRT
- <213> Homo Sapien

agacettete eteetgeaaa tag 823

- <400> 132
- Met Tyr Arg His Lys Asn Ser Trp Arg Leu Gly Leu Lys Tyr Pro 1 5 10 15
- Pro Ser Ser Lys Glu Glu Thr Gln Val Pro Lys Thr Leu Ile Ser 20 25 30
- Gly Leu Pro Gly Arg Lys Ser Ser Ser Arg Val Gly Glu Lys Leu.

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Gln Ser Ala His Lys Met Pro Leu Ser Pro Gly Leu Leu Leu
                   50
                                       55
                                                            60
 Leu Leu Ser Gly Ala Thr Ala Thr Ala Ala Leu Pro Leu Glu Gly
 Gly Pro Thr Gly Arg Asp Ser Glu His Met Gln Glu Ala Ala Gly
                  80
                                       85
 Ile Arg Lys Ser Ser Leu Leu Thr Phe Leu Ala Trp Trp Phe Glu
                  95
                                      100
                                                          105
 Trp Thr Ser Gln Ala Ser Ala Gly Pro Leu Ile Gly Glu Glu Ala
                 110
                                      115
                                                          120
 Arg Glu Val Ala Arg Arg Gln Glu Gly Ala Pro Pro Gln Gln Ser
                 125
                                      130
                                                          135
 Ala Arg Arg Asp Arg Met Pro Cys Arg Asn Phe Phe Trp Lys Thr
                                      145
                 140
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 Phe Ser Ser Cys Lys
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<210> 133
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide probe
<400> 133
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<210> 134
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<220>
<223> Synthetic oligonucleotide probe
<400> 134
gcaggaggag aaggtcttcc agaagaag 28
<210> 135
<211> 45
<212> DNA
<213> Artificial Sequence
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<223> Synthetic oligonucleotide probe
<400> 135
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<210> 136
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- <211> 1875
- <212> DNA
- <213> Homo Sapien
- <400> 136 gtcgtgtgct tggaggaagc cgcggaaccc ccagcgtccg tccatggcgt 50 ggagccttgg gagctggctg ggtggctgcc tgctggtgtc agcattggga 100 atggtaccac ctcccgaaaa tgtcagaatg aattctgtta atttcaagaa 150 cattetacag tgggagtcac etgettttge caaagggaac etgaetttca 200 cagctcagta cctaagttat aggatattcc aagataaatg catgaatact 250 accttgacgg aatgtgattt ctcaagtctt tccaagtatg gtgaccacac 300 cttgagagtc agggctgaat ttgcagatga gcattcagac tgggtaaaca 350 tcaccttctg tcctgtggat gacaccatta ttggaccccc tggaatgcaa 400 gtagaagtac ttgctgattc tttacatatg cgtttcttag cccctaaaat 450 tgagaatgaa tacgaaactt ggactatgaa gaatgtgtat aactcatgga 500 cttataatgt gcaatactgg aaaaacggta ctgatgaaaa gtttcaaatt 550 actccccagt atgactttga ggtcctcaga aacctggagc catggacaac 600 ttattgtgtt caagttcgag ggtttcttcc tgatcggaac aaagctgggg 650 aatggagtga gcctgtctgt gagcaaacaa cccatgacga aacggtcccc 700 tcctggatgg tggccgtcat cctcatggcc tcggtcttca tggtctgcct 750 ggcactcctc ggctgcttct ccttgctgtg gtgcgtttac aagaagacaa 800 agtacgcctt ctcccctagg aattctcttc cacagcacct gaaagagttt 850 ttgggccatc ctcatcataa cacacttctg tttttctcct ttccattgtc 900 ggatgagaat gatgtttttg acaagctaag tgtcattgca gaagactctg 950 agageggeaa geagaateet ggtgaeaget geageetegg gaeeeegeet 1000 gggcagggc cccaaagcta ggctctgaga aggaaacaca ctcggctggg 1050 cacagtgacg tactccatct cacatctgcc tcagtgaggg atcagggcag 1100 caaacaaggg ccaagaccat ctgagccagc cccacatcta gaactccaga 1150 cctggactta gccaccagag agctacattt taaaggctgt cttggcaaaa 1200 atactccatt tgggaactca ctgccttata aaggctttca tgatgttttc 1250 agaagttggc cactgagagt gtaattttca gccttttata tcactaaaat 1300 aagatcatgt tttaattgtg agaaacaggg ccgagcacag tggctcacgc 1350

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<210> 137

<211> 325

<212> PRT

<213> Homo Sapien

<400> 137

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Ser Ala Leu Gly Met Val Pro Pro Pro Glu Asn Val Arg Met Asn 20 25 30

Ser Val Asn Phe Lys Asn Ile Leu Gln Trp Glu Ser Pro Ala Phe 35 40 45

Ala Lys Gly Asn Leu Thr Phe Thr Ala Gln Tyr Leu Ser Tyr Arg
50 55 60

Ile Phe Gln Asp Lys Cys Met Asn Thr Thr Leu Thr Glu Cys Asp
65 70 75

Phe Ser Ser Leu Ser Lys Tyr Gly Asp His Thr Leu Arg Val Arg 80 85 90

Ala Glu Phe Ala Asp Glu His Ser Asp Trp Val Asn Ile Thr Phe 95 100 105

Cys Pro Val Asp Asp Thr Ile Ile Gly Pro Pro Gly Met Gln Val

Glu Val Leu Ala Asp Ser Leu His Met Arg Phe Leu Ala Pro Lys 125 130 135

Ile Glu Asn Glu Tyr Glu Thr Trp Thr Met Lys Asn Val Tyr Asn 140 145 150

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Ser Trp Thr Tyr Asn Val Gln Tyr Trp Lys Asn Gly Thr Asp Glu
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                                      160
                                                          165
Lys Phe Gln Ile Thr Pro Gln Tyr Asp Phe Glu Val Leu Arg Asn
                 170
                                      175
                                                           180
Leu Glu Pro Trp Thr Thr Tyr Cys Val Gln Val Arg Gly Phe Leu
                 185
                                      190
                                                          195
Pro Asp Arg Asn Lys Ala Gly Glu Trp Ser Glu Pro Val Cys Glu
                 200
                                      205
                                                          210
Gln Thr Thr His Asp Glu Thr Val Pro Ser Trp Met Val Ala Val
                 215
                                      220
                                                          225
Ile Leu Met Ala Ser Val Phe Met Val Cys Leu Ala Leu Leu Gly
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                                     235
                                                          240
Cys Phe Ser Leu Leu Trp Cys Val Tyr Lys Lys Thr Lys Tyr Ala
                 245
                                     250
                                                          255
Phe Ser Pro Arg Asn Ser Leu Pro Gln His Leu Lys Glu Phe Leu
                 260
                                     265
                                                          270
Gly His Pro His His Asn Thr Leu Leu Phe Phe Ser Phe Pro Leu
                 275
                                     280
                                                          285
Ser Asp Glu Asn Asp Val Phe Asp Lys Leu Ser Val Ile Ala Glu
                 290
                                     295
                                                          300
Asp Ser Glu Ser Gly Lys Gln Asn Pro Gly Asp Ser Cys Ser Leu
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Gly Thr Pro Pro Gly Gln Gly Pro Gln Ser
                 320
                                     325
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- <210> 138
- <211> 2570
- <212> DNA
- <213> Homo Sapien
- <400> 138
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 geegeegeeg etgetggegg agatgeeeeg eegggeaaaa tegeggtggt 150
 tggggetggg attggggget etgetgtgge eeattttete eageageaet 200
 ttggaeeteg ggtgeagate gaegtgtaeg agaagggaae egtgggtgge 250
 egettggeea eeateteagt eaacaageag eactatgaga geggggetge 300
 eteetteeae teeetgagee tgeacatgea ggaettegte aagetgetgg 350
 ggetgaggea eeggeggag gtggtgggea ggagegeeat etteggegg 400

gagcacttca tgctggagga gactgactgg tacctgctga acctcttccg 450 cctctggtgg cactatggca tcagcttcct gaggctgcag atgtgggtgg 500 aggaggtcat ggagaagttc atgaggatct ataagtacca ggcccacggc 550 tatgccttct cgggtgtgga ggagctgctc tactcactgg gggagtccac 600 ctttgttaac atgacccagc actctgtggc tgagtccctg ctgcaggtgg 650 gcgtcacgca gcgctttatt gatgatgtcg tttctgctgt cctgcgggcc 700 agctatggcc agtcagcagc gatgcccgcc tttgcaggag ccatgtcact 750 agccggggcc caaggcagcc tgtggtctgt ggaaggaggc aataagctgg 800 tttgttccgg tttgctgaag ctcaccaagg ccaatgtgat ccatgccaca 850 gtgacctctg tgaccctgca cagcacagag gggaaagccc tgtaccaggt 900 ggcgtatgag aatgaggtag gcaacagctc tgacttctat gacatcgtgg 950 tcatcgccac cccctgcac ctggacaaca gcagcagcaa cttaaccttt 1000 graggetter accepted tgatgacgtg raggetett teragerear 1050 cgtcgtctcc ttggtccacg gctacctcaa ctcgtcctac ttcggtttcc 1100 cagaccctaa gcttttcccc tttgccaaca tccttaccac agatttcccc 1150 agettettet geactetgga caacatetge eetgteaaca tetetgeeag 1200 cttccggcga aagcagcccc aggaggcagc tgtttggcga gtccagtccc 1250 ccaagcccct ctttcggacc cagctaaaga ccctgttccg ttcctattac 1300 tcagtgcaga cagctgagtg gcaggcccat ccctctatg gctcccgccc 1350 cacgctcccg aggtttgcac tccatgacca gctcttctac ctcaatgccc 1400 tggagtgggc ggccagctcc gtggaggtga tggccgtggc tgccaagaat 1450 gtggccttgc tggcttacaa ccgctggtac caggacctag acaagattga 1500 tcaaaaagat ttgatgcaca aggtcaagac tgaactgtga gggctctagg 1550 gagageetgg gaacttteat eecceatga agatggatea teecacagea 1600 gcccaggact gaataagcca tgctcgccca ccaggcttct ttctgacccc 1650 tcatgtatca agcatctcca ggtgacctac tgtctgccta tattaagggt 1700 ccacacggcg gctgctgctt ttttttaagg gggaaagtaa gaaaagagaa 1750 ggaaatccaa gccagtatat ttgttttatt tattttttt aagaagaaaa 1800 aagttcatct tcacaaggtg cttcagactt ggtttcttag ctagaaacca 1850

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<210> 139

<211> 494

<212> PRT

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Val Val Gly Ala Gly Ile Gly Gly Ser Ala Val Ala His Phe Leu 35 40 45

Gln Gln His Phe Gly Pro Arg Val Gln Ile Asp Val Tyr Glu Lys
50 55 60

Gly Thr Val Gly Gly Arg Leu Ala Thr Ile Ser Val Asn Lys Gln
65 70 75

His Tyr Glu Ser Gly Ala Ala Ser Phe His Ser Leu Ser Leu His 80 85 90

Met Gln Asp Phe Val Lys Leu Leu Gly Leu Arg His Arg Arg Glu 95 100 105

Val Val Gly Arg Ser Ala Ile Phe Gly Glu His Phe Met Leu

				110					115					120
Glu	Glu	Thr	Asp	Trp 125	Tyr	Leu	Leu	Asn	Leu 130	Phe	Arg	Leu	Trp	Trp 135
His	Tyr	Gly	Ile	Ser 140	Phe	Leu	Arg	Leu	Gln 145	Met	Trp	Val	Glu	Glu 150
Val	Met	Glu	Lys	Phe 155	Met	Arg	Ile	Tyr	Lys 160	Tyr	Gln	Ala	His	Gly 165
Tyr	Ala	Phe	Ser	Gly 170	Val	Glu	Glu	Leu	Leu 175	Tyr	Ser	Leu	Gly	Glu 180
Ser	Thr	Phe	Val	Asn 185	Met	Thr	Gln	His	Ser 190	Val	Ala	Glu	Ser	Leu 195
Leu	Gln	Val	Gly	Val 200	Thr	Gln	Arg	Phe	Ile 205	Asp	Asp	Val	Val	Ser 210
Ala	Val	Leu	Arg	Ala 215	Ser	Tyr	Gly	Gln	Ser 220	Ala	Ala	Met	Pro	Ala 225
Phe	Ala	Gly	Ala	Met 230	Ser	Leu	Ala	Gly	Ala 235	Gln	Gly	Ser	Leu	Trp 240
Ser	Val	Glu	Gly	Gly 245	Asn	Lys	Leụ	Val	Cys 250	Ser	Gly	Leu	Leu	Lys 255
Leu	Thr	Lys	Ala	Asn 260	Val	Ile	His	Ala	Thr 265	Val	Thr	Ser	Val	Thr 270
Leu	His	Ser	Thr	Glu 275	Gly	Lys	Ala	Leu	Tyr 280	Gln	Val	Ala	Tyr	Glu 285
Asn	Glu	Val	Gly	Asn 290	Ser	Ser	Asp	Phe	Tyr 295	Asp	Ile	Val	Val	Ile 300
Ala	Thr	Pro	Leu	His 305	Leu	Asp	Asn	Ser	Ser 310	Ser	Asn	Leu	Thr	Phe 315
Ala	Gly	Phe	His	Pro 320	Pro	Ile	Asp	Asp	Val 325	Gln	Gly	Ser	Phe	Gln 330
Pro	Thr	Val	Val	Ser 335	Leu	Val	His	Gly	Tyr 340	Leu	Asn	Ser	Ser	Tyr 345
Phe	Gly	Phe	Pro	Asp 350	Pro	Lys	Leu	Phe	Pro 355	Phe	Ala	Asn	Ile	Leu 360
Thr	Thr	Asp	Phe	Pro 365	Ser	Phe	Phe	Cys	Thr 370	Leu	Asp	Asn	Ile	Cys 375
Pro	Val	Asn	,Ile	Ser 380	Ala	Ser	Phe	Arg	Arg 385	Lys	Gln	Pro	Gln	Glu 390
Ala	Ala	Val	Trp	Arg 395	Val	Gln	Ser	Pro	Lys 400	Pro	Leu	Phe	Arg	Thr 405

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Glu Trp Gln Ala His Pro Leu Tyr Gly Ser Arg Pro Thr Leu Pro
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                 425
                                      430
Arg Phe Ala Leu His Asp Gln Leu Phe Tyr Leu Asn Ala Leu Glu
                                                           450
                                      445
                 440
 Trp Ala Ala Ser Ser Val Glu Val Met Ala Val Ala Ala Lys Asn
                                                           465
                 455
                                      460
Val Ala Leu Leu Ala Tyr Asn Arg Trp Tyr Gln Asp Leu Asp Lys
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